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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/202,549D

DATE: 08/21/2002

TIME: 15:19:43

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\08212002\I202549D.raw

4 <110> APPLICANT: Tsichlis, Philip  
 5 Grimes, Leighton H  
 6 Zweidler-McKay, Patrick  
 9 <120> TITLE OF INVENTION: NUCLEIC ACID MOLECULE FOR ENHANCING GENE EXPRESSION  
 12 <130> FILE REFERENCE: FCCC96-11  
 15 <140> CURRENT APPLICATION NUMBER: US 09/202,549D  
 16 <141> CURRENT FILING DATE: 1999-10-12  
 19 <150> PRIOR APPLICATION NUMBER: PCT/US97/10486  
 20 <151> PRIOR FILING DATE: 1997-06-17  
 23 <150> PRIOR APPLICATION NUMBER: US 60/019,808  
 24 <151> PRIOR FILING DATE: 1996-06-17  
 27 <160> NUMBER OF SEQ ID NOS: 70  
 30 <170> SOFTWARE: PatentIn version 3.1  
 33 <210> SEQ ID NO: 1  
 34 <211> LENGTH: 12  
 35 <212> TYPE: DNA  
 36 <213> ORGANISM: Artificial Sequence  
 W--> 37 <220> FEATURE:  
 38 <223> OTHER INFORMATION: Gfi-1 binding sequence  
 W--> 39 <220> FEATURE:  
 40 <221> NAME/KEY: misc\_feature  
 41 <222> LOCATION: (1)..(1)  
 42 <223> OTHER INFORMATION: "n" is any nucleotide  
 W--> 44 <220> FEATURE:  
 45 <221> NAME/KEY: misc\_feature  
 46 <222> LOCATION: (9)..(9)  
 47 <223> OTHER INFORMATION: "n" is any nucleotide  
 49 <400> SEQUENCE: 1  
 W--> 50 naaatcacng ca 12  
 53 <210> SEQ ID NO: 2  
 54 <211> LENGTH: 12  
 55 <212> TYPE: DNA  
 56 <213> ORGANISM: Artificial Sequence  
 58 <220> FEATURE:  
 59 <223> OTHER INFORMATION: Gfi-1 binding sequence  
 W--> 60 <220> FEATURE:  
 61 <221> NAME/KEY: misc\_feature  
 62 <222> LOCATION: (9)..(9)  
 63 <223> OTHER INFORMATION: "n" is "t" or "a"  
 65 <400> SEQUENCE: 2  
 W--> 66 taaatcacng ca 12  
 69 <210> SEQ ID NO: 3  
 70 <211> LENGTH: 12

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Input Set : A:\PTO.VSK.txt  
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71 <212> TYPE: DNA  
72 <213> ORGANISM: Artificial Sequence  
74 <220> FEATURE:  
75 <223> OTHER INFORMATION: An expression regulatory DNA segment  
W--> 76 <220> FEATURE:  
77 <221> NAME/KEY: misc\_feature  
78 <222> LOCATION: (1)..(1)  
79 <223> OTHER INFORMATION: "n" is any nucleotide  
W--> 81 <220> FEATURE:  
82 <221> NAME/KEY: misc\_feature  
83 <222> LOCATION: (3)..(3)  
84 <223> OTHER INFORMATION: "n" is any nucleotide  
W--> 86 <220> FEATURE:  
87 <221> NAME/KEY: misc\_feature  
88 <222> LOCATION: (4)..(5)  
89 <223> OTHER INFORMATION: "n" is "g" "c" or "t", or is absent, or is an oligonucleotide  
of  
90 two or more nucleotides  
W--> 92 <220> FEATURE:  
93 <221> NAME/KEY: misc\_feature  
94 <222> LOCATION: (6)..(6)  
95 <223> OTHER INFORMATION: "n" is "a" "g" or "c", or is absent, or is an oligonucleotide  
of  
96 two or more nucleotides  
W--> 98 <220> FEATURE:  
99 <221> NAME/KEY: misc\_feature  
100 <222> LOCATION: (9)..(9)  
101 <223> OTHER INFORMATION: "n" is "a" "g" or "c", or is absent, or is an  
oligonucleotide of  
102 two or more nucleotides  
104 <400> SEQUENCE: 3  
W--> 105 nannnnacng ca 12  
108 <210> SEQ ID NO: 4  
109 <211> LENGTH: 24  
110 <212> TYPE: DNA  
111 <213> ORGANISM: Artificial Sequence  
113 <220> FEATURE:  
114 <223> OTHER INFORMATION: Gfi-1 binding sequence  
W--> 115 <220> FEATURE:  
116 <221> NAME/KEY: misc\_feature  
117 <222> LOCATION: (2)..(2)  
118 <223> OTHER INFORMATION: "n" is "a" or "c"  
W--> 121 <220> FEATURE:  
122 <221> NAME/KEY: misc\_feature  
123 <222> LOCATION: (7)..(7)  
124 <223> OTHER INFORMATION: "n" is "inosine" or "c"  
W--> 126 <220> FEATURE:  
127 <221> NAME/KEY: misc\_feature  
128 <222> LOCATION: (15)..(15)  
129 <223> OTHER INFORMATION: "n" is "a" or "t"  
131 <400> SEQUENCE: 4  
W--> 132 anaaaanaaa tcacngcata tgcc 24

RAW SEQUENCE LISTING  
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Input Set : A:\PTO.VSK.txt  
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135 <210> SEQ ID NO: 5  
136 <211> LENGTH: 33  
137 <212> TYPE: DNA  
138 <213> ORGANISM: Artificial Sequence  
140 <220> FEATURE:  
141 <223> OTHER INFORMATION: Gfi-1 binding sequence  
143 <400> SEQUENCE: 5  
144 accatcacca cataaaatcac tgccttatcct gtg 33  
147 <210> SEQ ID NO: 6  
148 <211> LENGTH: 33  
149 <212> TYPE: DNA  
150 <213> ORGANISM: Artificial Sequence  
152 <220> FEATURE:  
153 <223> OTHER INFORMATION: Gfi-1 binding oligonucleotide  
155 <400> SEQUENCE: 6  
156 accatcacca cataaaatcac tgccttatcct gtg 33  
159 <210> SEQ ID NO: 7  
160 <211> LENGTH: 24  
161 <212> TYPE: DNA  
162 <213> ORGANISM: Artificial Sequence  
164 <220> FEATURE:  
165 <223> OTHER INFORMATION: Gfi-1 binding oligonucleotide  
167 <400> SEQUENCE: 7  
168 caccacatag atcactgcct atcc 24  
171 <210> SEQ ID NO: 8  
172 <211> LENGTH: 24  
173 <212> TYPE: DNA  
174 <213> ORGANISM: Artificial Sequence  
176 <220> FEATURE:  
177 <223> OTHER INFORMATION: Gfi-1 binding oligonucleotide  
179 <400> SEQUENCE: 8  
180 caccacataa ctcactgcct atcc 24  
183 <210> SEQ ID NO: 9  
184 <211> LENGTH: 24  
185 <212> TYPE: DNA  
186 <213> ORGANISM: Artificial Sequence  
188 <220> FEATURE:  
189 <223> OTHER INFORMATION: Gfi-1 binding oligonucleotide  
191 <400> SEQUENCE: 9  
192 caccacataa ataactgcct atcc 24  
195 <210> SEQ ID NO: 10  
196 <211> LENGTH: 24  
197 <212> TYPE: DNA  
198 <213> ORGANISM: Artificial Sequence  
200 <220> FEATURE:  
201 <223> OTHER INFORMATION: Gfi-1 binding oligonucleotide  
203 <400> SEQUENCE: 10  
204 caccacataa atcaatgcct atcc 24  
207 <210> SEQ ID NO: 11

RAW SEQUENCE LISTING  
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Input Set : A:\PTO.VSK.txt  
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208 <211> LENGTH: 24  
209 <212> TYPE: DNA  
210 <213> ORGANISM: Artificial Sequence  
212 <220> FEATURE:  
213 <223> OTHER INFORMATION: Gfi-1 binding oligonucleotide  
215 <400> SEQUENCE: 11  
216 caccacataa atcacttcct atcc 24  
219 <210> SEQ ID NO: 12  
220 <211> LENGTH: 500  
221 <212> TYPE: DNA  
222 <213> ORGANISM: Artificial Sequence  
224 <220> FEATURE:  
225 <223> OTHER INFORMATION: Gfi-1 binding sequence  
227 <400> SEQUENCE: 12  
228 gccccctgg ctgaccggcc aacgacccccc cgggattgac gtcaataatg acgtatgttc 60  
230 ccatagtaac gccaataggg actttccatt gacgtcaatg ggtggagat ttacqgtaaa 120  
232 ctgcccactt ggcagtacat caagtgtatc atatgccaag tacgccccctt attgacgtca 180  
234 atgacggtaa atggcccgcc tggcattatg cccagtgatc yaccilatgg gactttccct 240  
236 cttggcagta catctacgta tttagtcatcg ctattaccat ggtgatgcgg ttttggcagt 300  
238 acatcaatgg gcgtggatag cggtttgact cacggggatt tccaagtctc cacccattg 360  
240 acgtcaatgg gagtttggttt tggcacccaa atcaacggga ctttccaaaa tgtcgtaaca 420  
242 actccgcccc attgacgcaa atggccggta ggcgtgtacg gtgggaggtc tatataagca 480  
244 gagctcggtt agtgaaccgt 500  
247 <210> SEQ ID NO: 13  
248 <211> LENGTH: 500  
249 <212> TYPE: DNA  
250 <213> ORGANISM: Artificial Sequence  
252 <220> FEATURE:  
253 <223> OTHER INFORMATION: Gfi-1 binding sequence  
255 <400> SEQUENCE: 13  
256 gccccctgg ctgaccggcc aacgacccccc cgggattgac gtcaataatg acgtatgttc 60  
258 ccatagtaac gccaataggg actttccatt gacgtcaatg ggtggagat ttacqgtaaa 120  
260 ctgcccactt ggcagtacat caagtgtatc atatgccaag tacgccccctt attgacgtca 180  
262 atgacggtaa atggcccgcc tggcattatg cccagtgatc gaccttatgg gactttccct 240  
264 cttggcagta catctacgta tttagtcatcg ctattaccat ggtgatgcgg ttttggcagt 300  
266 acatcaatgg gcgtggatag cggtttgact cacggggagt tccaagtctc cacccattg 360  
268 acgtcaatgg gagtttggttt tggcacccaa ctcaacggga ctttccaaaa tgtcgtaaca 420  
270 actccgcccc attgacgcaa atggccggta ggcgtgtacg gtgggaggtc tatataagca 480  
272 gagctcggtt agtgaaccgt 500  
275 <210> SEQ ID NO: 14  
276 <211> LENGTH: 500  
277 <212> TYPE: DNA  
278 <213> ORGANISM: Artificial Sequence  
280 <220> FEATURE:  
281 <223> OTHER INFORMATION: Gfi-1 binding sequence  
283 <400> SEQUENCE: 14  
284 gccccctgg ctgaccggcc aacgacccccc cgggattgac gtcaataatg acgtatgttc 60  
286 ccatagtaac gccaataggg actttccatt gacgtcaatg ggtggagat ttacqgtaaa 120  
288 ctgcccactt ggcagtacat caagtgtatc atatgccaag tacgccccctt attgacgtca 180

RAW SEQUENCE LISTING  
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Input Set : A:\PTO.VSK.txt  
Output Set: N:\CRF4\08212002\I202549D.raw

290 atgacggtaa atggcccgcc tggcattatg cccagttacat gaccttatgg gactttccta 240  
292 cttggcagta catctacgta ttagtcatcg ctattaccat ggtgatgcgg tttggcagt 300  
294 acatcaatgg gcgtggatag cggtttgact cacgggactt tccaaagtctc cacccattg 360  
296 acgtcaatgg gagtttggtt tggcaccaaa actaacggga ctttccaaaa tgtcgtaaca 420  
298 actccgcccc attgacgcaa atggcggtt ggcgtgtacg gtgggaggtc tatataagca 480  
300 gagctcggtt agtgaaccgt 500  
303 <210> SEQ ID NO: 15  
304 <211> LENGTH: 12  
305 <212> TYPE: DNA  
306 <213> ORGANISM: Artificial Sequence  
308 <220> FEATURE:  
309 <223> OTHER INFORMATION: Gfi-1 binding sequence  
311 <400> SEQUENCE: 15  
312 caaatcaata ac 12  
315 <210> SEQ ID NO: 16  
316 <211> LENGTH: 12  
317 <212> TYPE: DNA  
318 <213> ORGANISM: Artificial Sequence  
320 <220> FEATURE:  
321 <223> OTHER INFORMATION: Gfi-1 binding sequence  
323 <400> SEQUENCE: 16  
324 taaatctgtg tg 12  
327 <210> SEQ ID NO: 17  
328 <211> LENGTH: 12  
329 <212> TYPE: DNA  
330 <213> ORGANISM: Artificial Sequence  
333 <220> FEATURE:  
334 <223> OTHER INFORMATION: Gfi-1 binding sequence  
336 <400> SEQUENCE: 17  
337 gaaatcagtt aa 12  
340 <210> SEQ ID NO: 18  
341 <211> LENGTH: 12  
342 <212> TYPE: DNA  
343 <213> ORGANISM: Artificial Sequence  
346 <220> FEATURE:  
347 <223> OTHER INFORMATION: Gfi-1 binding sequence  
349 <400> SEQUENCE: 18  
350 gaaatcagac ca 12  
353 <210> SEQ ID NO: 19  
354 <211> LENGTH: 12  
355 <212> TYPE: DNA  
356 <213> ORGANISM: Artificial Sequence  
359 <220> FEATURE:  
360 <223> OTHER INFORMATION: Gfi-1 binding sequence  
362 <400> SEQUENCE: 19  
363 gaaatcagtt aa 12  
366 <210> SEQ ID NO: 20  
367 <211> LENGTH: 12  
368 <212> TYPE: DNA

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/202,549D

DATE: 08/21/2002  
TIME: 15:19:45

Input Set : A:\PTO.VSK.txt  
Output Set: N:\CRF4\08212002\I202549D.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 1,9  
Seq#:2; N Pos. 9  
Seq#:3; N Pos. 1,3,4,5,6,9  
Seq#:4; N Pos. 2,7,15

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/202,549D

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TIME: 15:19:45

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\08212002\I202549D.raw

L:37 M:283 W: Missing Blank Line separator, <220> field identifier  
L:39 M:283 W: Missing Blank Line separator, <220> field identifier  
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:60 M:283 W: Missing Blank Line separator, <220> field identifier  
L:66 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:76 M:283 W: Missing Blank Line separator, <220> field identifier  
L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:115 M:283 W: Missing Blank Line separator, <220> field identifier  
L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0  
L:562 M:283 W: Missing Blank Line separator, <220> field identifier